

SEQUENCE LISTING

<110> Masure, Stefan
Richardson, Alan

<120> Human AKT-3

<130> 51869/001

<140> PCT/GB99/04311

<141> 1999-12-17

<150> GB 9828375.7

<151> 1998-12-22

<160> 3

<170> PatentIn Ver. 2.0

<210> 1

<211> 1547

<212> DNA

<213> Homo sapiens

<400> 1

gggagtcatc atgagcgatg ttaccattgt gaaagaaggt tgggttcaga agaggggaga
60

atatataaaa aactggaggc caagatactt ctttttgaag acagatggct cattcatagg
120

atataaagag aaacctcaag atgtggattt accttatccc ctcaacaact tttcagtggc
180

aaaatgccag ttaatgaaaa cagaacgacc aaagccaaac acatttataa tcagatgtct
240

[illegible]

acgagaataa gtctctttca ttctgtact tcaactgtcat cttcaattta ttactgaaaa
1500

tgattcctgg acatcaccag tcctagctct tacacatagc aggggca
1547

<210> 2

<211> 1435

<212> DNA

<213> Homo sapiens

<400> 2

atgagc gatg ttaccattgt gaaagaaggt tgggttcaga agaggggaga atatataaaa
60

aactggaggc caagatactt ccttttgaag acagatggct cattcatagg atataaagag
120

aaacctcaag atgtggattt accttatccc ctcaacaact tttcagtggc aaaatgccag
180

ttaatgaaaa cagaacgacc aaagccaaac acatttataa tcagatgtct ccagtggact
240

actgttatag agagaacatt tcatgtagat actccagagg aaaggggaaga atggacagaa
300

gctatccagg ctgtagcaga cagactgcag aggcaagaag aggagagaat gaattgtagt
360

ccaacttcac aaattgataa tataggagag gaagagatgg atgcctctac aacccatcat
420

aaaagaaaga caatgaatga ttttgactat ttgaaactac taggtaaagg cacttttggg
480

aaagttattt tggttcgaga gaaggcaagt ggaaaatact atgctatgaa gattctgaag
540

aaagaagtca ttattgcaaa ggatgaagtg gcacacactc taactgaaag cagagtatta
600

aagaacacta gacatccctt ttttaacatcc ttgaaatatt ccttcagac aaaagaccgt
660

ttgtgttttg tgatggaata tgttaatggg ggcgagctgt tttccattt gtcgagagag
720

cgggtgttct ctgaggaccg cacacgttcc tatggtgcag aaattgtctc tgccttggac
780

tatctacatt ccggaaagat tgtgtaccgt gatctcaagt tggagaatct aatgctggac
840

aaagatggcc acataaaaat tacagatttt ggactttgca aagaagggat cacagatgca
900

000000-000000

gacaatgaga ggcggccgca tttccctcaa ttttcctact ctgcaagtga acgaga
1436

<213> Homo sapiens

1 5 10 15

20 25 30

35 40 45

Tyr Pro Leu Asn Asn Phe Ser Val Ala Lys Cys Gln Leu Met Lys Thr

60

80

95

110

125

140

160

175

190

205

220

Leu Ser Ser Asp Ala Lys Ser Leu Leu Ser Gly Leu Leu Ile Lys Asp
370 375 380

Pro Asn Lys Arg Leu Gly Gly Gly Pro Asp Asp Ala Lys Glu Ile Met
 385 390 395 400

Arg His Ser Phe Phe Ser Gly Val Asn Trp Gln Asp Val Tyr Asp Lys
 405 410 415

Lys Leu Val Pro Pro Phe Lys Pro Gln Val Thr Ser Glu Thr Asp Thr
 420 425 430

Arg Tyr Phe Asp Glu Glu Phe Thr Ala Gln Thr Ile Thr Ile Thr Pro
 435 440 445

Pro Glu Lys Tyr Asp Glu Asp Gly Met Asp Cys Met Asp Asn Glu Arg
 450 455 460

Arg Pro His Phe Pro Gln Phe Ser Tyr Ser Ala Ser Gly Arg Glu
 465 470 475

100290-6205550